

pAS2 Insert	Colony Color		
	pGAD GH Insert		
	PIN1	PIN2	PIN3
NIMA	Blue	Blue	Blue
K40M NIMA	Blue	Blue	Blue
NIMA280-699	Blue	Blue	White
NLK1	White	White	Blue
NIMA280-699FS	White	White	White

FIG. 1A

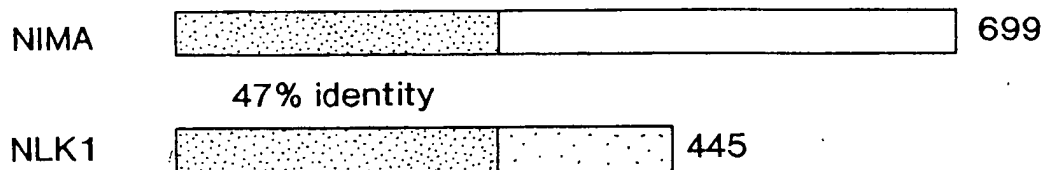


FIG. 1B

TGCTGGCCAGCACCTCGAGGGAAG

1 ATGGCGGACGAGGAGAAGCTGCCGCCCCGGCTGGGAGAAGCGCATGAGCCGAGCTCAGGC 20
M A D E E K L P P G W E K R M S R S S G
61 CGAGTGTAATACTTCAACCACATCACTAACGCCAGCCAGTGGGAGCGGCCCAGCGGCAAC 40
R V Y Y F N H I T N A S Q W E R P S G N
121 AGCAGCAGTGGTGGCAAAAACGGGCAGGGGGAGCCTGCCAGGGTCCGCTGCTCGCACCTG 60
S S S G G K N G Q G E P A R V R C S H L
181 CTGGTGAAGCACAGCCAGTCACGGCGGCCCTCGTCCTGGCGGCAGGAGAAGATCACCCGG 80
L V K H S Q S R R P S S W R Q E K I T R
241 ACCAAGGAGGAGGCCCTGGAGCTGATCAACGGCTACATCAGAAAGATCAAGTCGGGAGAG 100
T K E E A L E L I N G Y I Q K I K S G E
301 GAGGACTTTGAGTCTCTGGCCTCACAGTTCAGCGACTGCAGCTCAGCCAAGGCCAGGGGA 120
E D F E S L A S Q F S D C S S A K A R G
361 GACCTGGGTGCCTTCAGCAGAGGTCAGATGCAGAAGCCATTTGAAGACGCCTCGTTTGCG 140
D L G A F S R G Q M Q K P F E D A S F A
421 CTGCGGACGGGGGAGATGAGCGGGCCCGTGTTACGGATTCCGGCATCCACATCATCCTC 160
L R T G E M S G P V F T D S G I H I I L
481 CGCACTGAGTGAGGGTGGGGAGCCCAGGCCTGGCCTCGGGGCAGGGCAGGGCGGCTAGGC
R T E *
541 CGGCCAGCTCCCCCTTGCCCGCCAGCCAGTGGCCGAACCCCCACTCCCTGCCACCGTCA
601 CACAGTATTTATTGTTCACAAATGGCTGGGAGGGGGCCCTTCCAGATTGGGGGCCCTGG
661 GGTCCCCACTCCCTGTCCATCCCCAGTTGGGGCTGCGACCGCCAGATTCTCCCTTAAGGA
721 ATTGACTTCAGCAGGGGTGGGAGGCTCCCAGACCCAGGGCAGTGTGGTGGGAGGGGTGTT
781 CCAAAGAGAAGGCCTGGTCAGCAGAGCCGCCCCGTGTCCCCCAGGTGCTGGAGGCAGAC
841 TCGAGGGCCGAATTGTTTCTAGTTAGGCCACGCTCCTCTGTTACAGTCGAAAGGTGAACA
901 CTCATGCGGCAGCCATGGGCCCTCTGAGCAACTGTGCAGACCCTTTCACCCCCAATTAAA
961 CCCAGAACCACTAAAAAAAAAAAAAAAAAAAA

FIG. 2A

Pin1/Human	5	E K L P P G W E K R M S R S S G R V Y Y F N H I T N A S Q W E E R P S G N S S S
ESS1/SC	9	T G L P T P W T V R Y S K S K R E Y F F N P E T K H S Q W E E P E G T N K D
Yap/Human	171	V P L P A G W E M A K T S S - G Q R Y F L N H I D Q T T T W Q D P P R K A M L S
Nedd4/Mouse	?	S P L P P G W E E R Q D V L - G R T Y Y V V D H N T R T T Q W K R P S P D D D L
RSP5/SC	228	G R L P P G W E E R T D N F - G R T Y Y V V D H N T R T T Q W K R P S P D D D L
Drd/Human	3052	T S V Q G P W E R A I S P N K V P - Y Y I N H E T Q T T C W D H P P K M T E L Y
FE65/Rat	12	S D L P A G W M R V Q D T S - G - T Y Y W H I P T G T T Q W E P P G R A S P S
Consensus		. . L P . G W E G . . Y Y . N H . T . . T . W P

FIG. 2B

Pin1/Human	59	H L L V K H S Q S R R P S S W R Q E K I T R T K E E A L E L I N G Y I Q K I K
ESS1/SC	63	H I L I K H K D S R R P A S H R S E N I T I S K Q D A T D E L K T L I T R L D
Parvulin/EC	8	H I L V - - - - - - - - - - K E E K L A L - - D L L E Q I K
PrsA/BS	140	H I L V A D K K T - - - - - A E E V E K - - - - - K L K
Cbf2/CJ	137	H I L V A T - - - - - - - E K E A K D I I N E L K G L K
EST/AT	?	H I L V E I K
Consensus		S G E - - E D F E S L A S Q F S D C S S A K A - R G D L G A F S R - G Q M Q K
		D D S K T N S F E A L A K E R S D C S S Y K R - G G D L G W F F G R - G E M Q P
		K G E K - - - F E D L A K E Y S T D S S A S K - G G D L G W F F A K E G Q M D E
		G K E L D A K F S E L A K E K S I D P G S K N Q G G E L G W F - D Q S T M V K
		V S K - - A N F E E V A T R V S D C S S A K R - G G D L G S F G R - G Q M Q K
	 F E . L A K . . S . C S S . K . . G G D L G . F . R . G Q M . .
		P F E D A S F A L R T G E M S G - P V F T D S G I H I I L R T E 164 Pin1/Human
		S F E D A A F Q L K V G E V S D - I V E S G S G V H V I K R V G 170 ESS1/SC
		A F D K V V F C C P V L E P T G - P L H T Q Q F G Y H I I K V L Y 93 Parvulin/EC
		T F S K A A F K L K T G E V S D - P V K T Q Q F G Y H I I K K T E 215 PrsA/BS
		P F T D A A F A L K N G T I T T P V K T N F G Y H V I L K E N 219 Cbf2/CJ
		P F E E A T Y A L K V G D I S D - I V D T D S G V H I I K R T E 69 EST/AT
		. F . D A A F . L K . G E . S . . P V . T . . G Y H I I K Consensus

FIG. 2C

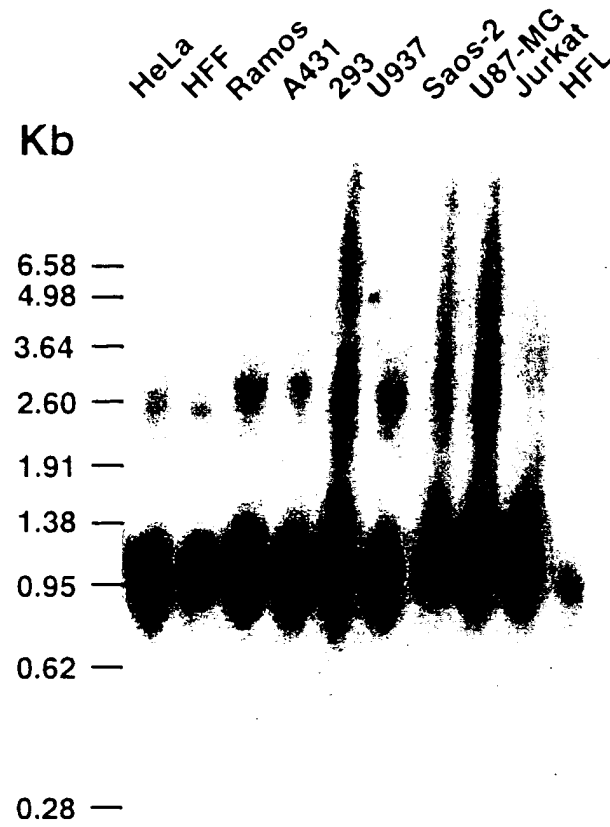
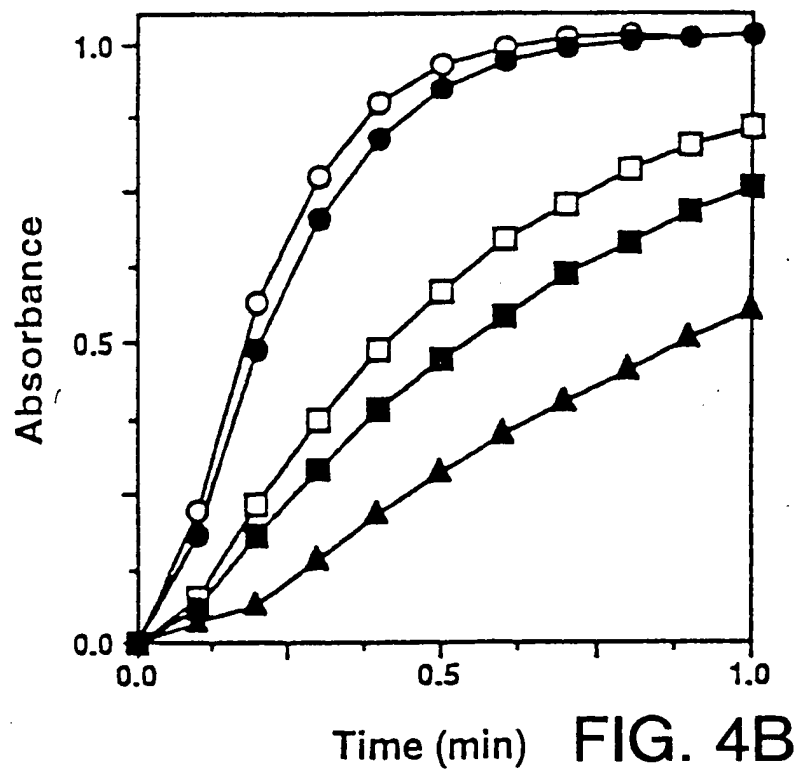
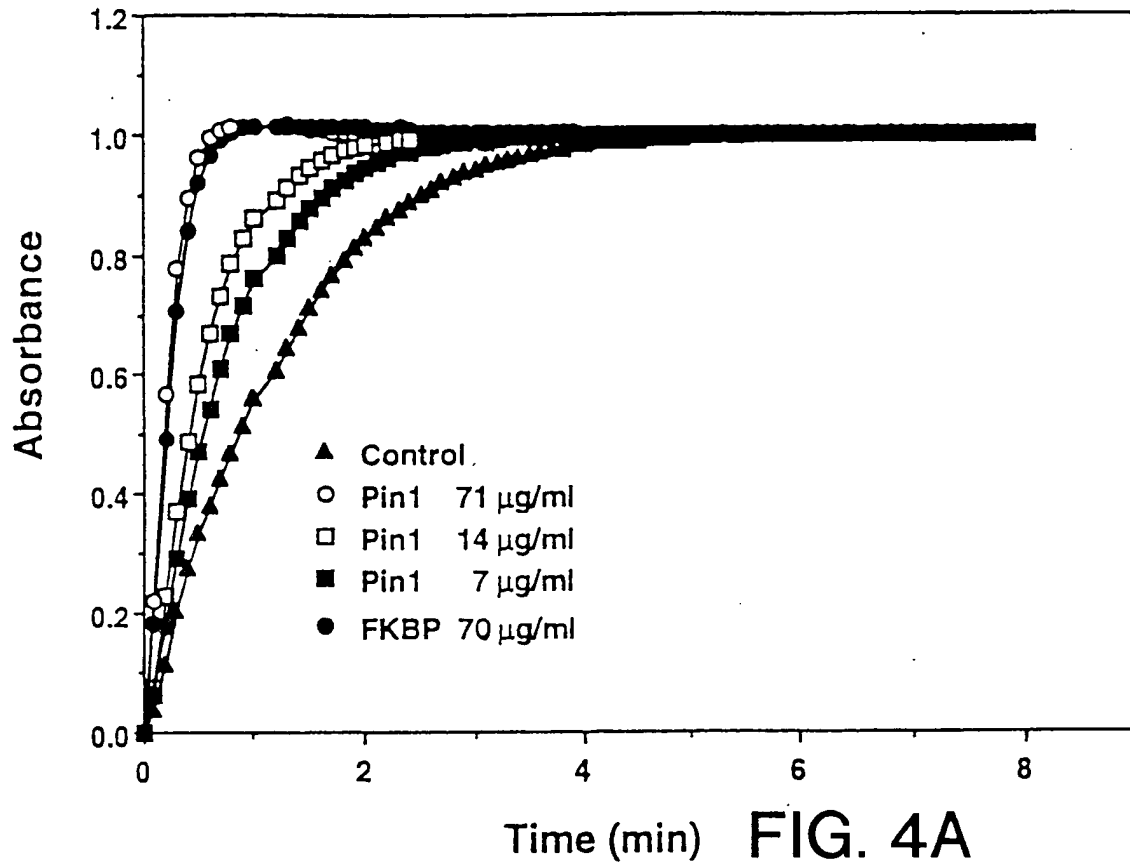
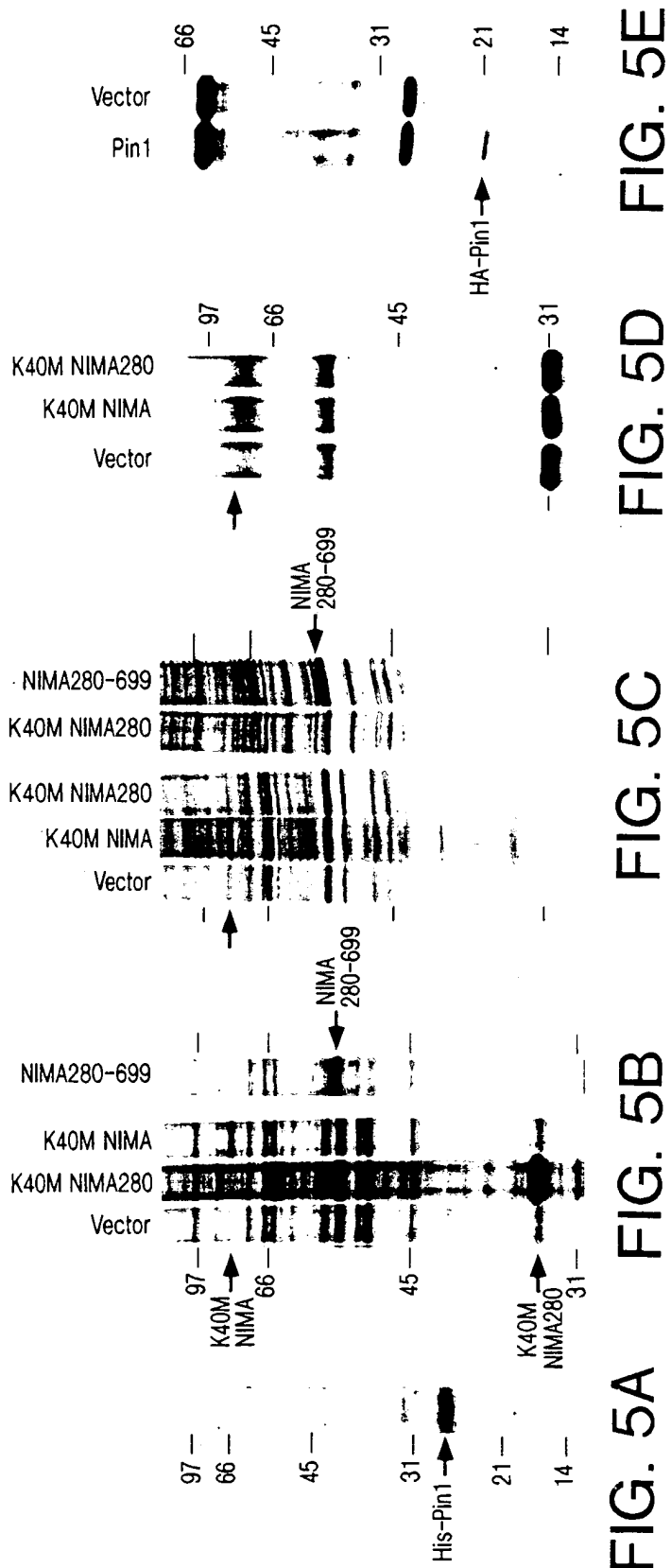


FIG. 3





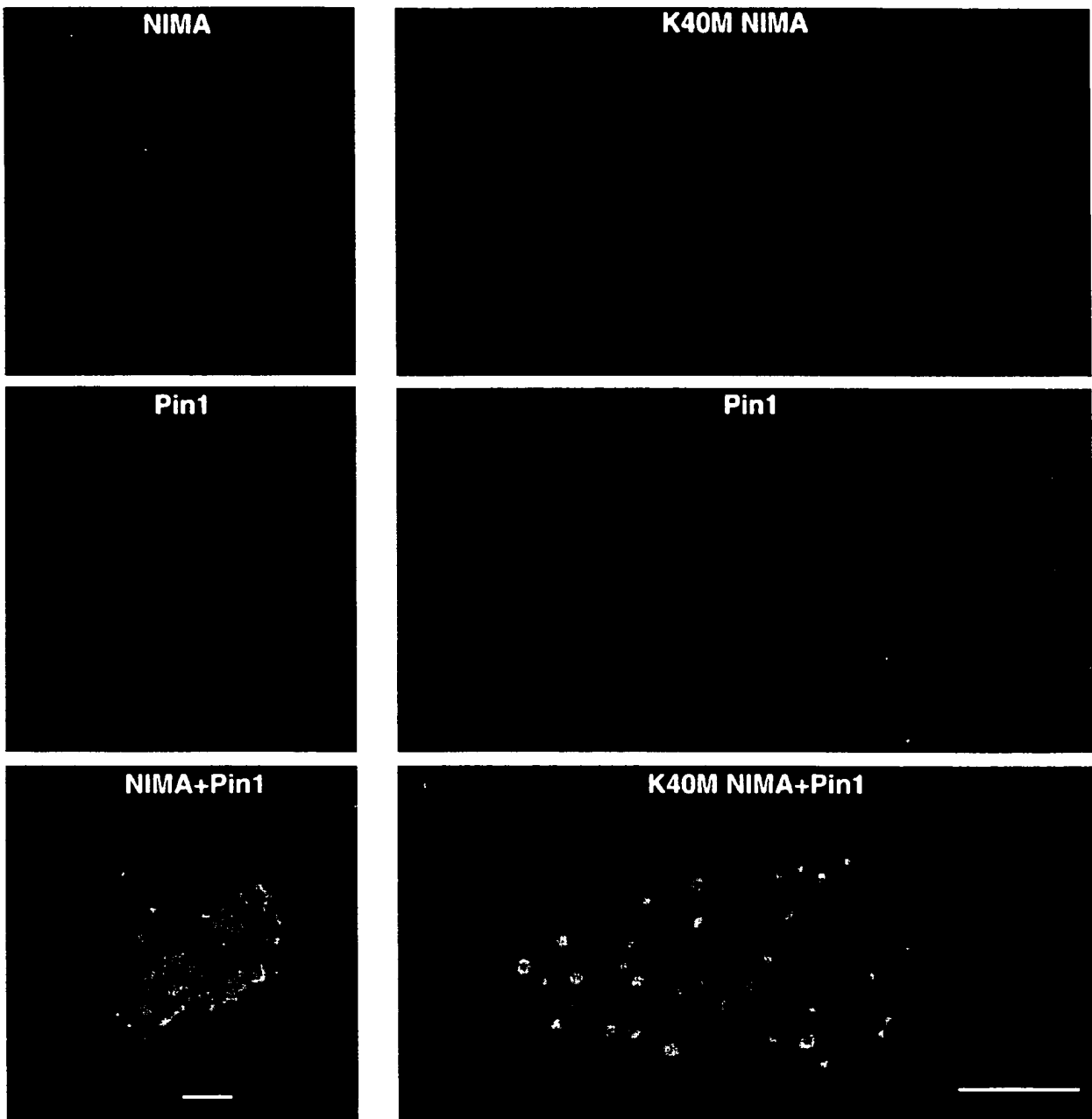


FIG. 6

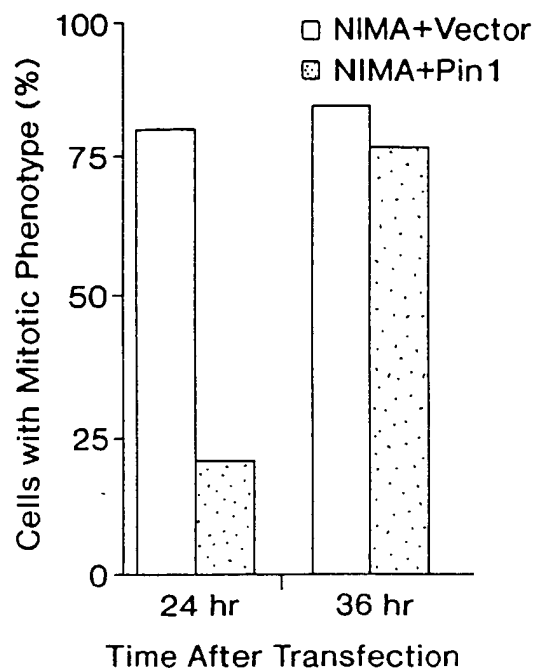


FIG. 7A

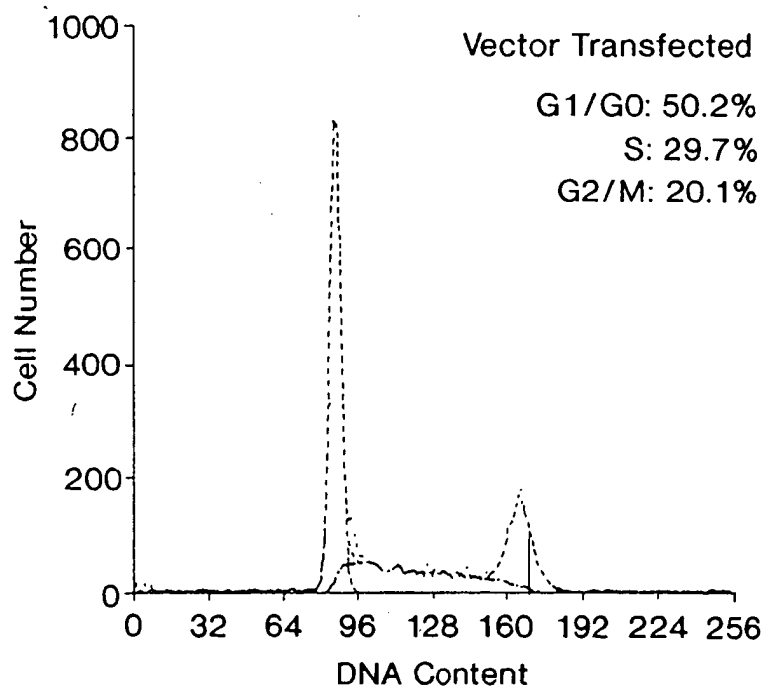


FIG. 7B

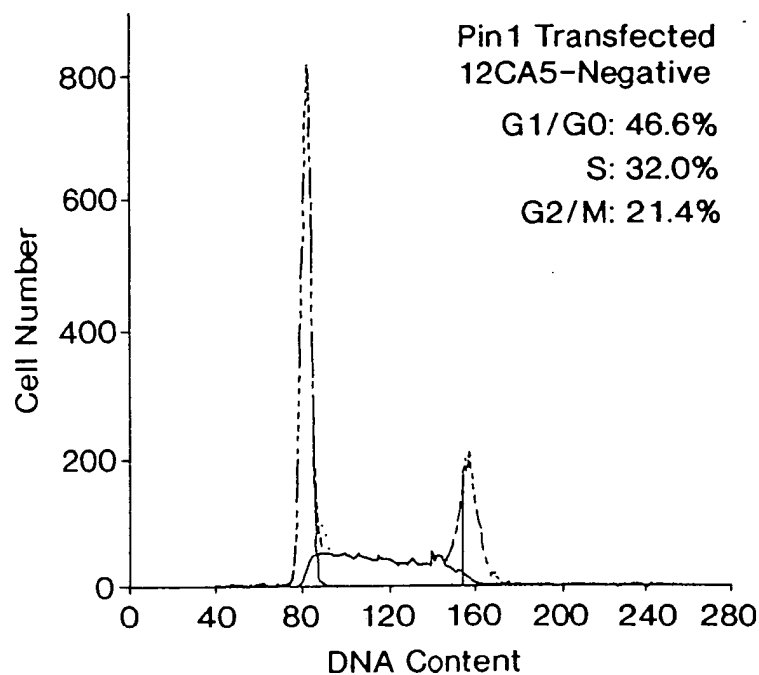


FIG. 7C

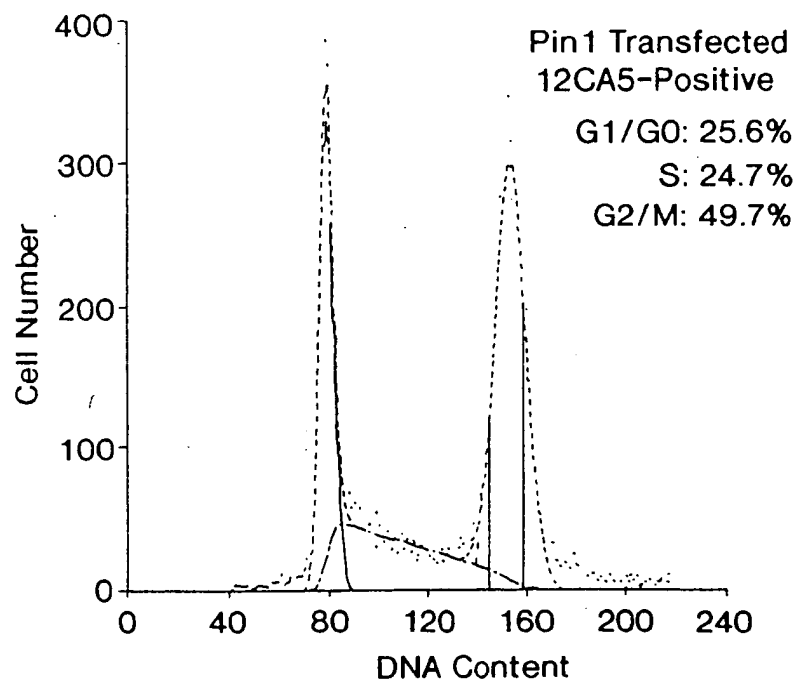


FIG. 7D

